

Jie Zhou
Building 12A, Room 4047
12 South Drive, MSC 5635
Bethesda, MD 20892
(301) 435-0378
zhoujie@mail.nih.gov

Summary of Qualification:

- Experienced Data management, statistical analysis, and pedigree data cleaning
- Experienced MS-access Database designing, coding, testing, supporting, and developing
- Experienced enhance database performance, perform restores, implement recovery procedures, regular system backups
- Experienced in developing and maintaining Sample tracking system
- Experienced in training understanding and using database system and sample tracking system
- Experienced and certified SAS programmer
- Experienced in other Statistical packages (R, SPSS), and Computer languages (Basic, C++, C, Java), and Genetic Packages (such as, Plink, PEDCHECK, Progeny etc.)
- Experienced in analyzing clinical, epidemiological, genetic, family base etc data, generating reports, tables, listings and graphs
- Experienced in extracting data and creating SAS datasets from various sources
- Experienced in developing, validating and implementing SAS programs and procedures derived datasets for analysis and generating & documenting tables, listings and graphs
- Experienced in using MS Access, Word, Excel, PowerPoint

Professional Experience:

2008 – Present: Data manager/ SAS programmer for Center Research Global Genomic Health (CRGGH)/NHGRI/NIH (NIH contractor from Kelly Service), Bethesda, Maryland

- Database designing, cording, testing, maintaining, and developing for CRGGH research studies
- Perform data QC checks to insure integrity and consistence of the dataset
- Maintaining, updating, backups and develop CRGGH sample tracking system
- Prepare and document properly data sets for researchers in CRGGH and collaborators
- Create, maintain and manage center datasets dictionaries and construction
- Provide and update Statistical analysis for various studies in CRGGH
- Provide genetic data analysis and create statistical reports, tables, listing and graphs to assist investigators in center and collaborators

2001 – 2008: Data manager/ SAS programmer for Division of Genetic Epidemiology, National Human Genome Center, School of Medicine, Howard University, Washington, D.C.

- Provided database design, maintain and develop for various studies in center
- Provided statistical programming support for center and collaborator.
- Assisted researches in center and collaborators for data analysis, statistical report interpretation, published tables, listing and graphs
- Designed and maintained sample tracking system for studies

Professional Education & Certification:

- 2000 – 2001: Computer Science, Loyola University at Chicago
- Nov.2003 SAS Certified Base Programmer
- Dec. 2001: S.A.G.E training course, Los Angeles CA
- May. 2002: SAS/INSIGHT training course, Rockville MD

- Nov. 2002: SAS/WEBTOOLS training course , Rockville MD
- Jun. 2008: Attend 2008 Summer Institute in Statistical Genetics, University of Washington
 - Module 3(Computing for Statistical Genetics);
 - Module 5 (Regression and Analysis of Variance);
 - Module 8: Quantitative Genetics;
 - Module 14: Human Association Mapping.
- Dec. 2008: SAS Enterprise Guide Overview training course, NIH
- May.2012: Scientific Graphics with Tips for PowerPoint, NIH
- Apr.2013: SPSS Tips & Tricks, NIH
- Jun 2013: PowerPoint Tips and Tricks, NIH
- Jun.2013: Introduction to Microsoft Access 2010, NIH
- July.2013: Intermediate Access 2010, NIH
- Sep. 2013: Advanced Excel 2010: PivotTables & More, NIH
- Sep.2013: Excel Formulas for Scientists, NIH
- Feb.2014: An introduction to R, NIH
- Feb.2014: Advanced Access 2010/ VBA Basics, NIH
- July 2014: Introduction to Macros, NIH
- Oct.2014: JavaScript with jQuery, Montgomery College MD

Award:

- 2012: Distinguished Achievement Award from Kelly government solutions management Team
- 2009: Distinguished Performance Award from Kelly government solutions management Team

Selected Publications:

- 1 Ramos E, Doumatey A, Elkahloun AG, Shriner D, Huang H, Chen G, **Zhou J**, McLeod H, Adeyemo A, Rotimi CN. Pharmacogenomics, ancestry and clinical decision making for global populations. *Pharmacogenomics J.* 2014 Jun;14(3):217-22. doi: 10.1038/tpj.2013.24. Epub 2013 Jul 9.
- 2 Marshall PA, Adebamowo CA, Adeyemo AA, Ogundiran TO, Strenski T, **Zhou J**, Rotimi CN. Voluntary participation and comprehension of informed consent in a genetic epidemiological study of breast cancer in Nigeria. *BMC Med Ethics.* 2014 May 13;15:38. doi: 10.1186/1472-6939-15-38.
- 3 Doumatey AP, **Zhou J**, Adeyemo A, Rotimi C. High sensitivity C-reactive protein (Hs-CRP) remains highly stable in long-term archived human serum. *Clin Biochem.* 2014 Mar;47(4-5):315-8. doi: 10.1016/j.clinbiochem.2013.12.014. Epub 2013 Dec 27.
- 4 Bentley AR, Chen G, Shriner D, Doumatey AP, **Zhou J**, Huang H, Mullikin JC, Blakesley RW, Hansen NF, Bouffard GG, Cherukuri PF, Maskeri B, Young AC, Adeyemo A, Rotimi CN. Gene-based sequencing identifies lipid-influencing variants with ethnicity-specific effects in African Americans. *PLoS Genet.* 2014 Mar 6;10(3):e1004190. doi: 10.1371/journal.pgen.1004190. eCollection 2014 Mar.
- 5 Monda KL, Chen GK, Taylor KC, Palmer C, Edwards TL, Lange LA, Ng MC, Adeyemo AA, Allison MA, Bielak LF, Chen G, Graff M, Irvin MR, Rhie SK, Li G, Liu Y, Liu Y, Lu Y, Nalls MA, Sun YV, Wojczynski MK, Yanek LR, Aldrich MC, Ademola A, **Zhou J**, Rotimi CN, et al. A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. *Nat Genet.* 2013 Jun;45(6):690-6. doi: 10.1038/ng.2608. Epub 2013 Apr 14.
- 6 Liu CT, Monda KL, Taylor KC, Lange L, Demerath EW, Palmas W, Wojczynski MK, Ellis JC, Vitolins MZ, Liu S, Papanicolaou GJ, Irvin MR, Xue L, Griffin PJ, Nalls MA, Adeyemo A, Liu J, Li G, Ruiz-Narvaez EA, Chen WM, Chen F, Henderson BE, Millikan RC, Ambrosone CB, Strom SS, Guo X, Andrews JS, Sun YV, Mosley TH, Yanek LR, Shriner D, Haritunians T, Rotter JI, Speliotes EK, Smith M, Rosenberg L, Mychaleckyj J, Nayak U, Spruill I, Garvey WT, Pettaway C, Nyante S, Bandera EV, Britton AF, Zonderman AB, Rasmussen-Torvik LJ, Chen YD, Ding J, Lohman K, Kritchevsky SB, Zhao W, Peyser PA, Kardia SL, Kabagambe E, Broeckel U, Chen G, **Zhou J**, Wassertheil-Smoller S, Neuhouser ML, Rampersaud E, Psaty B, Kooperberg C, Manson JE, Kuller LH, Ochs-Balcom HM, Johnson KC, Sucheston L, Ordovas JM, Palmer JR, Haiman CA, McKnight B, Howard BV, Becker DM, Bielak LF, Liu Y, Allison MA, Grant SF, Burke GL, Patel SR, Schreiner PJ, Borecki IB, Evans MK, Taylor H, Sale MM, Howard V, Carlson CS, Rotimi CN, Cushman M, Harris TB, Reiner AP, Cupples LA, North KE, Fox CS. Genome-wide association of body fat distribution in African ancestry populations suggests new loci. *PLoS Genet.* 2013;9(8):e1003681. doi: 10.1371/journal.pgen.1003681. Epub 2013 Aug 15.

- 7 Chen G, Bentley A, Adeyemo A, Shriner D, **Zhou J**, Doumatey A, Huang H, Ramos E, Erdos M, Gerry N, Herbert A, Christman M, Rotimi C. Genome-wide association study identifies novel loci association with fasting insulin and insulin resistance in African Americans. *Hum Mol Genet*. 2012 Oct 15;21(20):4530-6. Epub 2012 Jul 12.
- 8 Adeyemo A, Bentley AR, Meilleur KG, Doumatey AP, Chen G, **Zhou J**, Shriner D, Huang H, Herbert A, Gerry NP, Christman MF, Rotimi CN. Transferability and fine mapping of genome-wide associated loci for lipids in African Americans. *BMC Med Genet*. 2012 Sep 21;13:88. doi: 10.1186/1471-2350-13-88.
- 9 Doumatey AP, Chen G, Tekola Ayele F, **Zhou J**, Erdos M, Shriner D, Huang H, Adeleye J, Balogun W, Fasanmade O, Johnson T, Oli J, Okafor G, Amoah A, Eghan BA, Agyenim-Boateng K, Acheampong J, Adebamowo C, Gerry NP, Christman MF, Adeyemo A, Rotimi CN. C-reactive protein (CRP) promoter polymorphisms influence circulating CRP levels in a genome-wide association study of African Americans. *Hum Mol Genet*. 2012 Jul 1;21(13):3063-72. doi: 10.1093/hmg/ddz133. Epub 2012 Apr 5.
- 10 Tekola Ayele F, Doumatey A, Huang H, **Zhou J**, Charles B, Erdos M, Adeleye J, Balogun W, Fasanmade O, Johnson T, Oli J, Okafor G, Amoah A, Eghan BA Jr, Agyenim-Boateng K, Acheampong J, Adebamowo CA, Herbert A, Gerry N, Christman M, Chen G, Shriner D, Adeyemo A, Rotimi CN. Genome-wide associated loci influencing interleukin (IL)-10, IL-1Ra, and IL-6 levels in African Americans. *Immunogenetics*. 2012 May;64(5):351-9. doi: 10.1007/s00251-011-0596-7. Epub 2011 Dec 29.
- 11 Chen G, Ramos E, Adeyemo A, Shriner D, **Zhou J**, Doumatey AP, Huang H, Erdos MR, Gerry NP, Herbert A, Bentley AR, Xu H, Charles BA, Christman MF, Rotimi CN. UGT1A1 is a major locus influencing bilirubin levels in African Americans. *Eur J Hum Genet*. 2012 Apr;20(4):463-8. doi: 10.1038/ejhg.2011.206. Epub 2011 Nov 16.
- 12 Doumatey AP, Bentley AR, **Zhou J**, Huang H, Adeyemo A, Rotimi CN. Paradoxical Hyperadiponectinemia is Associated With the Metabolically Healthy Obese (MHO) Phenotype in African Americans. *J Endocrinol Metab*. 2012 Apr 1;2(2):51-65.
- 13 Chen G, Yuan A, Zhou Y, Bentley AR, **Zhou J**, Chen W, Shriner D, Adeyemo A, Rotimi CN. Simultaneous Analysis of Common and Rare Variants in Complex Traits: Application to SNPs (SCARVAAsnp). *Bioinform Biol Insights*. 2012;6:177-85. doi: 10.4137/BBI.S9966. Epub 2012 Jul 31.
- 14 Chen G, Yuan A, Zhou Y, Bentley AR, **Zhou J**, Chen W, Shriner D, Adeyemo A, Rotimi CN. Simultaneous Analysis of Common and Rare Variants in Complex Traits: Application to SNPs (SCARVAAsnp). *Bioinform Biol Insights*. 2012;6:177-85. doi: 10.4137/BBI.S9966. Epub 2012 Jul 31.
- 15 Bentley AR, Doumatey AP, Chen G, Huang H, **Zhou J**, Shriner D, Jiang C, Zhang Z, Liu G, Fasanmade O, Johnson T, Oli J, Okafor G, Eghan BA Jr, Agyenim-Boateng K, Adeleye J, Balogun W, Adebamowo C, Amoah A, Acheampong J, Adeyemo A, Rotimi CN. Simple f test reveals gene-gene interactions in case-control studies. *Int J Nephrol*. 2012;2012:748984. doi: 10.1155/2012/748984. Epub 2012 Sep 2.
- 16 Doumatey AP, **Zhou J**, Huang H, Adeleye J, Balogun W, Fasanmade O, Johnson T, Oli J, Okafor G, Amoah A, Eghan B, Agyenim-Boateng K, Acheampong J, Adebamowo C, Adeyemo A, Rotimi CN. Circulating adiponectin is associated with renal function independent of age and serum lipids in west africans. *Int J Nephrol*. 2012;2012:730920. doi: 10.1155/2012/730920. Epub 2012 Aug 22.
- 17 Shriner D, Herbert A, Doumatey AP, **Zhou J**, Huang H, Erdos MR, Chen G, Gerry NP, Christman MF, Adeyemo A, Rotimi CN. Multiple loci associated with renal function in African Americans. *PLoS One*. 2012;7(9):e45112. doi: 10.1371/journal.pone.0045112. Epub 2012 Sep 13.
- 18 Charles BA, Doumatey A, Huang H, **Zhou J**, Chen G, Shriner D, Adeyemo A, Rotimi CN. The roles of IL-6, IL-10, and IL-1RA in obesity and insulin resistance in African-Americans. *J Clin Endocrinol Metab*. 2011 Dec;96(12):E2018-22. doi: 10.1210/jc.2011-1497. Epub 2011 Sep 28.
- 19 Faruque MU, Chen G, Doumatey A, Huang H, **Zhou J**, Dunston GM, Rotimi CN, Adeyemo AA. Association of ATP1B1, RG55 and SELE polymorphisms with hypertension and blood pressure in African-Americans. *J Hypertens*. 2011 Oct;29(10):1906-12. doi: 10.1097/HJH.0b013e32834b000d.
- 20 Ramos E, Chen G, Shriner D, Doumatey A, Gerry NP, Herbert A, Huang H, **Zhou J**, Christman MF, Adeyemo A, Rotimi C. Replication of genome-wide association studies (GWAS) loci for fasting plasma glucose in African-Americans. *Diabetologia*. 2011 Apr;54(4):783-8. doi: 10.1007/s00125-010-2002-7. Epub 2010 Dec 25.
- 21 Charles BA, Shriner D, Doumatey A, Chen G, **Zhou J**, Huang H, Herbert A, Gerry NP, Christman MF, Adeyemo A, Rotimi CN. A genome-wide association study of serum uric acid in African Americans. *BMC Med Genomics*. 2011 Feb 4;4:17. doi: 10.1186/1755-8794-4-17.
- 22 Sumner AE, **Zhou J**, Doumatey A, Imoisili OE, Amoah A, Acheampong J, Oli J, Johnson T, Adebamowo C, Rotimi CN. Low HDL-Cholesterol with Normal Triglyceride Levels is the Most Common Lipid Pattern in West Africans and African Americans with Metabolic Syndrome: Implications for Cardiovascular Disease Prevention. *CVD Prev Control*. 2010 Sep 1;5(3):75-80.

- 23 Chen G, Shriner D, **Zhou J**, Doumatey A, Huang H, Gerry NP, Herbert A, Christman MF, Chen Y, Dunston GM, Faruque MU, Rotimi CN, Adeyemo A. Development of admixture mapping panels for African Americans from commercial high-density SNP arrays. *BMC Genomics*. 2010 Jul 5;11:417. doi: 10.1186/1471-2164-11-417.
- 24 Meilleur KG, Doumatey A, Huang H, Charles B, Chen G, **Zhou J**, Shriner D, Adeyemo A, Rotimi C. Circulating adiponectin is associated with obesity and serum lipids in West Africans. *J Clin Endocrinol Metab*. 2010 Jul;95(7):3517-21. doi: 10.1210/jc.2009-2765. Epub 2010 Apr 9.
- 25 Adeyemo A, Chen G, **Zhou J**, Shriner D, Doumatey A, Huang H, Rotimi C. FTO genetic variation and association with obesity in West Africans and African Americans. *Diabetes*. 2010 Jun;59(6):1549-54. doi: 10.2337/db09-1252. Epub 2010 Mar 18.
- 26 Doumatey AP, Lashley KS, Huang H, **Zhou J**, Chen G, Amoah A, Agyenim-Boateng K, Oli J, Fasanmade O, Adebamowo CA, Adeyemo AA, Rotimi CN. Relationships among obesity, inflammation, and insulin resistance in African Americans and West Africans. *Obesity (Silver Spring)*. 2010 Mar;18(3):598-603. doi: 10.1038/oby.2009.322. Epub 2009 Oct 1.
- 27 Shriner D, Adeyemo A, Gerry NP, Herbert A, Chen G, Doumatey A, Huang H, **Zhou J**, Christman MF, Rotimi CN. Transferability and fine-mapping of genome-wide associated loci for adult height across human populations. *PLoS One*. 2009 Dec 22;4(12):e8398. doi: 10.1371/journal.pone.0008398.
- 28 Adeyemo A, Gerry N, Chen G, Herbert A, Doumatey A, Huang H, **Zhou J**, Lashley K, Chen Y, Christman M, Rotimi C. A genome-wide association study of hypertension and blood pressure in African Americans. *PLoS Genet*. 2009 Jul;5(7):e1000564. doi: 10.1371/journal.pgen.1000564. Epub 2009 Jul 17.
- 29 Chen G, Adeyemo A, **Zhou J**, Chen Y, Huang H, Doumatey A, Lashley K, Agyenim-Boateng K, Eghan BA Jr, Acheampong J, Fasanmade O, Johnson T, Okafor G, Oli J, Amoah A, Rotimi C. Genome-wide search for susceptibility genes to type 2 diabetes in West Africans: potential role of C-peptide. *Diabetes Res Clin Pract*. 2007 Dec;78(3):e1-6. Epub 2007 Jun 4.
- 30 Chen Y, Huang H, **Zhou J**, Doumatey A, Lashley K, Chen G, Agyenim-Boateng K, Eghan BA, Acheampong J, Fasanmade O, Johnson T, Akinsola FB, Okafor G, Oli J, Ezepue F, Amoah A, Akafo S, Adeyemo A, Rotimi CN. Polymorphism of the endothelial nitric oxide synthase gene is associated with diabetic retinopathy in a cohort of West Africans. *Mol Vis*. 2007 Nov 26;13:2142-7.
- 31 Gudmundsson J, Sulem P, Steinthorsdottir V, Bergthorsson JT, Thorleifsson G, Manolescu A, Rafnar T, Gudbjartsson D, Agnarsson BA, Baker A, Sigurdsson A, Benediktsdottir KR, Jakobsdottir M, Blöndal T, Stacey SN, Helgason A, Gunnarsdottir S, Olafsdottir A, Kristinsson KT, Birgisdottir B, Ghosh S, Thorlacius S, Magnusdottir D, Stefansdottir G, Kristjansson K, Bagger Y, Wilensky RL, Reilly MP, Morris AD, Kimber CH, Adeyemo A, Chen Y, **Zhou J**, So WY, Tong PC, Ng MC, Hansen T, Andersen G, Borch-Johnsen K, Jorgensen T, Tres A, Fuertes F, Ruiz-Echarri M, Asin L, Saez B, van Boven E, Klaver S, Swinkels DW, Aben KK, Graif T, Cashy J, Suarez BK, van Vierssen Trip O, Frigge ML, Ober C, Hofker MH, Wijmenga C, Christiansen C, Rader DJ, Palmer CN, Rotimi C, Chan JC, et al. Two variants on chromosome 17 confer prostate cancer risk, and the one in TCF2 protects against type 2 diabetes. *Nat Genet*. 2007 Aug;39(8):977-83. Epub 2007 Jul 1.
- 32 Steinthorsdottir V, Thorleifsson G, Reynisdottir I, Benediktsson R, Jonsdottir T, Walters GB, Styrkarsdottir U, Gretarsdottir S, Emilsson V, Ghosh S, Baker A, Snorradsdottir S, Bjarnason H, Ng MC, Hansen T, Bagger Y, Wilensky RL, Reilly MP, Adeyemo A, Chen Y, **Zhou J**, Gudnason V, Chen G, Huang H, Lashley K, Doumatey A, So WY, Ma RC, Andersen G, Borch-Johnsen K, Jorgensen T, van Vliet-Ostaptchouk JV, Hofker MH, Wijmenga C, Christiansen C, Rader DJ, Rotimi C, et al. A variant in CDKAL1 influences insulin response and risk of type 2 diabetes. *Nat Genet*. 2007 Jun;39(6):770-5. Epub 2007 Apr 26.
- 33 Chen G, Adeyemo AA, **Zhou J**, Chen Y, Doumatey A, Lashley K, Huang H, Amoah A, Agyenim-Boateng K, Eghan BA Jr, Okafor G, Acheampong J, Oli J, Fasanmade O, Johnson T, Rotimi C. A genome-wide search for linkage to renal function phenotypes in West Africans with type 2 diabetes. *Am J Kidney Dis*. 2007 Mar;49(3):394-400.
- 34 Helgason A, Pálsson S, Thorleifsson G, Grant SF, Emilsson V, Gunnarsdottir S, Adeyemo A, Chen Y, Chen G, Reynisdottir I, Benediktsson R, Hinney A, Hansen T, Andersen G, Borch-Johnsen K, Jorgensen T, Schäfer H, Faruque M, Doumatey A, **Zhou J**, Wilensky RL, Reilly MP, Rader DJ, Bagger Y, Christiansen C, Sigurdsson G, Hebebrand J, Pedersen O, Thorsteinsdottir U, Gulcher JR, Kong A, Rotimi C, Stefánsson K. Refining the impact of TCF7L2 gene variants on type 2 diabetes and adaptive evolution. *Nat Genet*. 2007 Feb;39(2):218-25. Epub 2007 Jan 7.
- 35 Marshall PA, Adebamowo CA, Adeyemo AA, Ogundiran TO, Vekich M, Strenski T, **Zhou J**, Prewitt TE, Cooper RS, Rotimi CN. Voluntary participation and informed consent to international genetic research. *Am J Public Health*. 2006 Nov;96(11):1989-95. Epub 2006 Oct 3.

- 36 Rotimi CN, Chen G, Adeyemo AA, Jones LS, Agyenim-Boateng K, Eghan BA Jr, **Zhou J**, Doumatey A, Lashley K, Huang H, Fasanmade O, Akinsola FB, Ezepue F, Amoah A, Akafu S, Chen Y, Oli J, Johnson T. Genomewide scan and fine mapping of quantitative trait loci for intraocular pressure on 5q and 14q in West Africans. *Invest Ophthalmol Vis Sci.* 2006 Aug;47(8):3262-7.
- 37 Chen G, Adeyemo A, **Zhou J**, Yuan A, Chen Y, Rotimi C. Genome scan linkage analysis comparing microsatellites and single-nucleotide polymorphisms markers for two measures of alcoholism in chromosomes 1, 4, and 7. *BMC Genet.* 2005 Dec 30;6 Suppl 1:S4.
- 38 Wilson JG, Rotimi CN, Ekunwe L, Royal CD, Crump ME, Wyatt SB, Steffes MW, Adeyemo A, **Zhou J**, Taylor HA Jr, Jaquish C. Study design for genetic analysis in the Jackson Heart Study. *Ethn Dis.* 2005 Autumn;15(4 Suppl 6):S6-30-37.
- 39 Adeyemo AA, Johnson T, Acheampong J, Oli J, Okafor G, Amoah A, Owusu S, Agyenim-Boateng K, Eghan BA Jr, Abbiyesku F, Fasanmade O, Rufus T, Doumatey A, Chen G, **Zhou J**, Chen Y, Furber-Harris P, Dunston G, Collins F, Rotimi C. A genome wide quantitative trait linkage analysis for serum lipids in type 2 diabetes in an African population. *Atherosclerosis.* 2005 Aug;181(2):389-97. Epub 2005 Feb 24.
- 40 Chen G, Adeyemo AA, Johnson T, **Zhou J**, Amoah A, Owusu S, Acheampong J, Agyenim-Boateng K, Eghan BA, Oli J, Okafor G, Abbiyesku F, Dunston GM, Chen Y, Collins F, Rotimi C. A genome-wide scan for quantitative trait loci linked to obesity phenotypes among West Africans. *Int J Obes (Lond).* 2005 Mar;29(3):255-9.
- 41 Chen Y, Kittles R, **Zhou J**, Chen G, Adeyemo A, Panguluri RK, Chen W, Amoah A, Opoku V, Acheampong J, Agyenim-Boateng K, Eghan BA Jr, Nyantaki A, Oli J, Okafor G, Ofoegbu E, Osotimehin B, Abbiyesku F, Johnson T, Fasanmade O, Rufus T, Furber-Harris P, Daniel HI, Berg KA, Collins FS, Dunston GM, Rotimi CN. Calpain-10 gene polymorphisms and type 2 diabetes in West Africans: the Africa America Diabetes Mellitus (AADM) Study. *Ann Epidemiol.* 2005 Feb;15(2):153-9.
- 42 Rotimi CN, Chen G, Adeyemo AA, Furber-Harris P, Parish-Gause D, **Zhou J**, Berg K, Adegoke O, Amoah A, Owusu S, Acheampong J, Agyenim-Boateng K, Eghan BA Jr, Oli J, Okafor G, Ofoegbu E, Osotimehin B, Abbiyesku F, Johnson T, Rufus T, Fasanmade O, Kittles R, Daniel H, Chen Y, Dunston G, Collins FS; Africa America Diabetes Mellitus (AADM) Study. A genome-wide search for type 2 diabetes susceptibility genes in West Africans: the Africa America Diabetes Mellitus (AADM) Study. *Diabetes.* 2004 Mar;53(3):838-41. Erratum in: *Diabetes.* 2004 May;53(5):1404. Guass, Debra [corrected to Parish-Gause, Debra].
- 43 Rotimi C, Daniel H, **Zhou J**, Obisesan A, Chen G, Chen Y, Amoah A, Opoku V, Acheampong J, Agyenim-Boateng K, Eghan BA Jr, Oli J, Okafor G, Ofoegbu E, Osotimehin B, Abbiyesku F, Johnson T, Fasanmade O, Doumatey A, Aje T, Collins F, Dunston G. Prevalence and determinants of diabetic retinopathy and cataracts in West African type 2 diabetes patients. *Ethn Dis.* 2003 Summer;13(2 Suppl 2):S110-7.